TTW AAG AAA ATC TCA TCC TCC F K K I S S S ် ၁၅၁ ၂ GCC AAC C GTC TCA GCA
V S A ည ပ ang ang con con tigg han ang ant tig ata M K R R R W K K N F I GCGCCGCCACC

ø

Gly/Ser-linker

SBP tag (SB19C4)

CGG GGC GGG GAG GAG GAG GGG GTG GGC R G G G AM WAY, W B G G B 299 TCC GGC GGC TGC AAG CTG S G G C K L GGC GAG CTG. GAG CTG. GGG GGG GTG. GAG CAC |CAC CCT. CAG GGG GAG GGG GGG G E E E E O L R A R E E H H P. O G G O R E P AGC GGT ACC ATG GAG GAG AGG ACG ACG GGG TGG S G S (M. D. E. K. T. T. G. 1900) GGTG CTT GGA AGC 666 6CA 6

b

* GCCCGGGCGGATCC

tag length: 78AA

2

G/S-linker

atelencians ace nes see testes see see see see sha site sas see see steres sas see see see see see AGC GGT ACC

လ ტ ß GGTG CTC GAG GGA AGC ß

GGC TCC S ტ TGC AAG CTG × ပ 299 G TCC GGC (S P. C.

AAG CGA EGA TGG AAA AAG AAT TTGAATANGCG GTG TCA EGA EGC RAGTCEG TUT AAG AAN ATC TGR TCC TGC GGG GCA CTT K R K K K T S S G A A A A A B K K T S S G A A

GGA

ტ

TAG GGCCCGAC

Tag length: 79 AA

17

Sequence

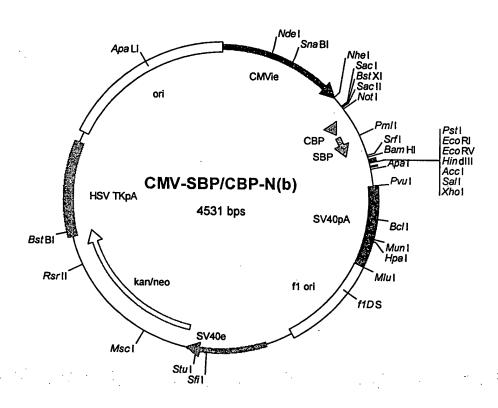
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581 583 583 584 585 585 587 589 5810 5811 5813	SB14 SB15 SB16 SB17 SB18 SB20

The "No." indicates the number of times each sequence was observed. The HPQ sequence is in bold type. Defined sequences at the termini are underlined. The six C-terminal residues are not shown.

Fig. &

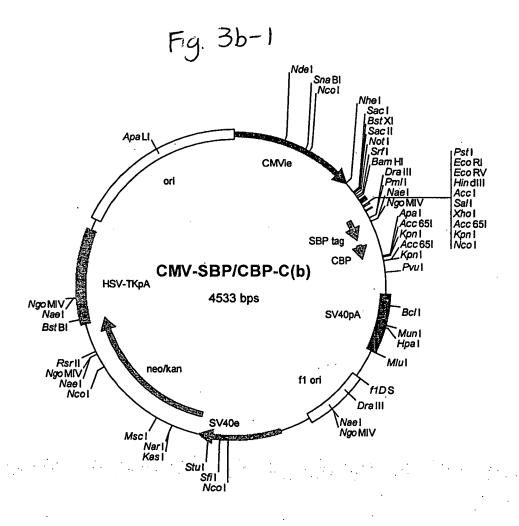
Fig. 3a-1



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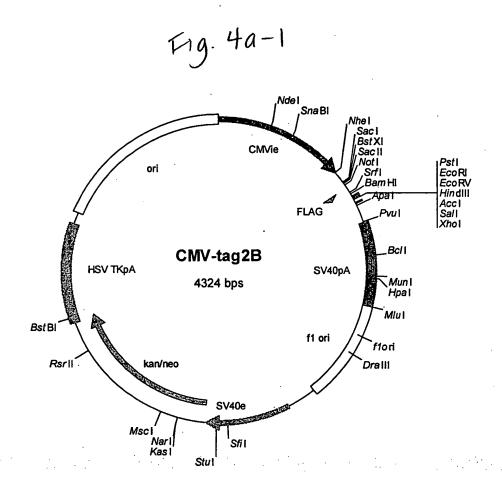


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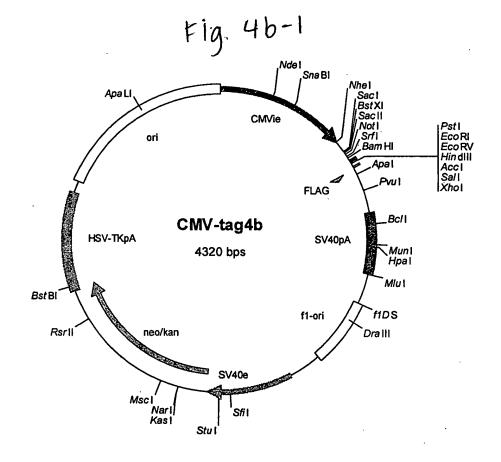
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Fig. 3b-3



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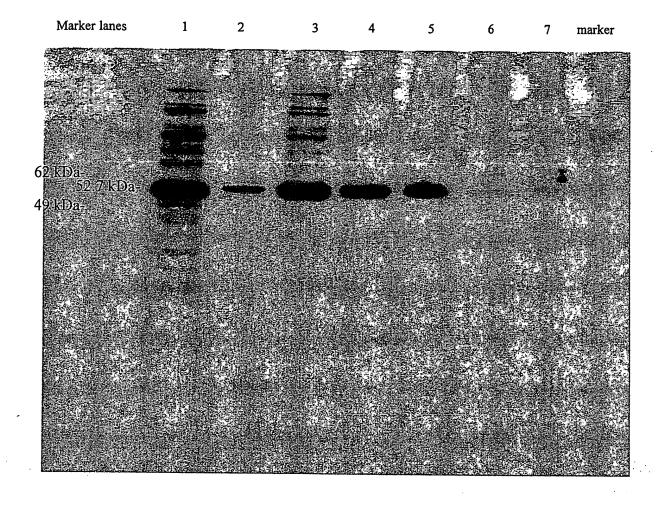
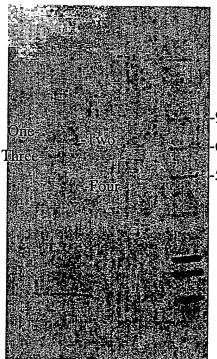


FIGURE 5



-98 kDa -64 kDa -50 kDa

FIGURE 6